

<110> Ferrick, David A. Swift, Susan E. Armstrong, Randall Fox, Bryan



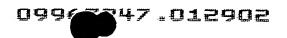
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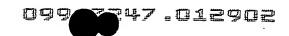
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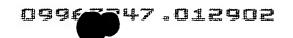
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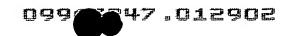
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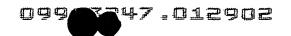


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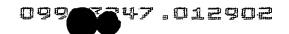
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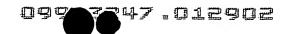
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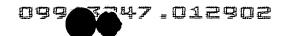


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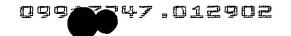
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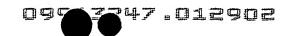
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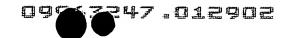
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Ala Val Lys Arg Pro Ala Ala Thr Lys Lys Ala Gly Gln Ala Lys Lys
                                   10
Lys Lys Leu Asp
           20
<210> 17
<211> 10
<212> PRT
<213> Artificial sequence
<220>
<223> stability sequence
<220>
<221> MISC_FEATURE
<222> (3)..(6)
<223> "Xaa" at positions 3 to 6 can be any amino acid.
<400> 17
Met Gly Xaa Xaa Xaa Gly Gly Pro Pro
               5
<210> 18
<211> 5
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Gly Ser Gly Gly Ser
<210> 19
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> linker sequence
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<400> 19
Gly Gly Gly Ser